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In the Claims

- 1. (amended) A compound that specifically inhibits the formation of the hu
 human C5b-9 complex selected from the group consisting of molecules structurally
 mimicking CD59 amino acid residues 42 to 58 when they are in a spatial orientation
 which inhibits formation of the hu C5b 9 complex, wherein the compound is not hu
 CD59 a peptidomimetic having the structure and function of human CD59 amino acid
 residues 42-58 of SEO ID NO:3 selected from the group consisting of a peptide, a nucleic
 acids acid, and a small molecule and an anti-ID anti-CD59 amino acid residues 42-58
 antibody, the peptidomimetic and an anti-CD59 antibody binding specifically to human
 C9 at amino acid residues 26-51 of SEO ID NO:14.
- 2. (amended) The compound of claim 1, selected from the group consisting of proteins, peptides, nucleic acids, and small molecules which bind specifically to amino acids 359 to 384 26-51 of hu human C9 in SEO ID NO:14.
- 3. (amended) The compound of claim 2, wherein the protein compound is an antibody.
- 4. (amended) The compound of claim 2, wherein the protein compound is a chimeric peptide which includes the amino acids 42 to 58 of the human sequence of CD59 in SEO ID NO:3.
- 5. (amended) The compound of claim 2, wherein the peptide compound is a covalently cyclized peptide comprising hu human CD59 amino acid residues 42 to 58 in SEO ID NO:3.

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- (amended) The compound of claim 2, wherein the composition compound 6. is a peptide of less than forty armino acids residues including amino acid residues 42 to 58 of hu human CD59 in SEQ ID NO:3
- (amended) The compound of claim-1, further A composition comprising a 7. compound that specifically inhibits the formation of the human C5b-9 complex selected from the group consisting of a peptidomimetic having the structure and function of human CD59 amino acid residues 42-58 of SEQ ID NO:3 selected from the group consisting of a peptide, a nucleic acid, and a small molecule and an anti-ID anti-CD59 amino acid residues 42-58 antibody, the peptidomimetic and an anti-CD59 antibody binding specifically to amino acid residues 26 to 51 of human C9 in SEQ ID NO:14, and a pharmaceutically acceptable carrier for administration to patients in need thercof.
- The compound of claim 1 wherein the compound is a peptidomimetic 8. compound comprising the side chains of hu human CD59 amino acid residues His44, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEO ID NO:3 in an equivalent spatial orientation and alignment to that presented on the surface of hu human CD59.
- The compound of claim 8 wherein the spatial orientation and alignment of 9. the side chains of His⁴⁴, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in the compound are equivalent to the spatial orientation and alignment deduced by NMR structure determination.
- (three times amended) A method for inhibiting human C5b-9 complex assembly comprising administering to a patient in need thereof an effective amount of a composition comprising a compound binding specifically to amino acid residues 26 to 51

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of human C9 in SEO ID NO:14 selected from the group consisting of a peptidomimetic having the structure and function of human CD59 amino acid residues 42-58 in SEO ID NO:3 selected from the group consisting of proteins, peptides a peptide, a nucleic acids acid, and a small molecule molecules having the structure and function of human CD59 amino acid residues 42-58, and an anti-ID anti-CD59 amino acid residues 42-58 antibody, the peptidomimetic and anti-CD59 antibody binding specifically to amino acid residues 359 384 of human C9.

- (three times amended) The method of claim 10, wherein the compound is 11. a peptidomimetic that is a small molecule which binds specifically to amino acids 359 to 384 26 to 51 of human C9 SEQ ID NO:14.
- (twice amended) The method of claim 10, wherein the protein compound 12. is an antibody.
- (three times amended) The method of claim 10, wherein the protein 13. compound is a chimeric peptide which includes the amino acids 42 to 58 of the human sequence of CD59 in SEQ ID NO:3.
- (three times amended) The method of claim 10, wherein the peptide compound is a covalently cyclized peptide comprising human CD59 amino acid residues 42 to 58 in SEQ ID NO:3.
- (three times amended) The method of claim 10, wherein the 15. peptidomimetic compound is a peptide of less than forty amino acids residues including amino acid residues 42 to 58 of human CD59 in SEQ ID NO:3.

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- (original) The method of claim 10, wherein the composition further 16. comprises a pharmaceutically acceptable carrier for administration to patients in need thereof.
- (once amended) The method of claim 10, wherein the composition is 17. administered to a patient is in need of suppression of complement-mediated inflammation.
- (twice amended) The method of claim 10 wherein the compound is a 18. peptidomimetic comprises comprising the side chains of human CD59 amino acid residues His⁴⁴, Asn⁴⁸, Asp⁴⁹, Thr⁵¹, Thr⁵², Arg⁵⁵, and Glu⁵⁸ of SEQ ID NO:3 in the spatial orientation and alignment of hu human CD59.
- (once amended) The method of claim 18 wherein the spatial orientation 19. and alignment of the side chains of His44, Asn48, Asp49, Thr51, Thr52, Arg55, and Glu58 of SEQ ID NO:3 in the compound are deduced by NMR structure determination.

Please cancel claims 20-35.